

FIG. 1A

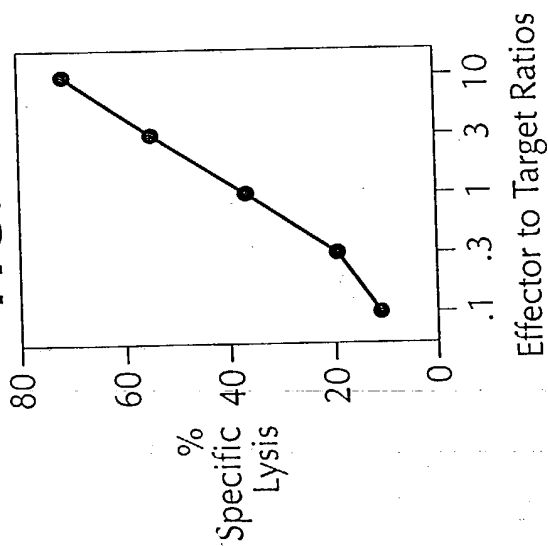


FIG. 1B

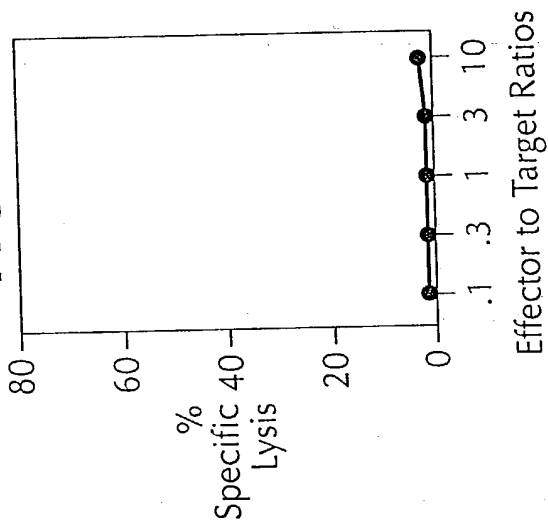


FIG. 1D

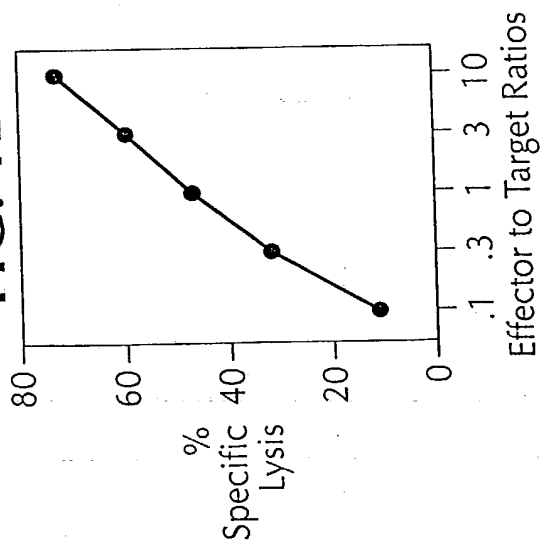


FIG. 1E

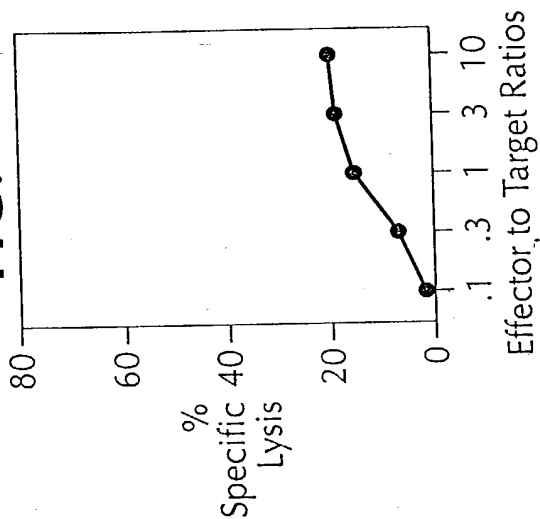


FIG. 1F

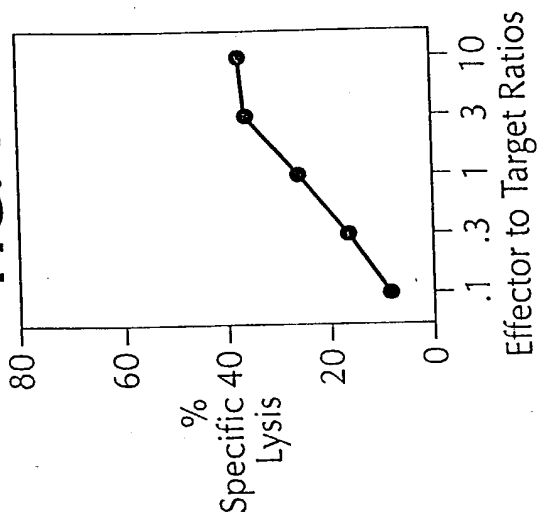
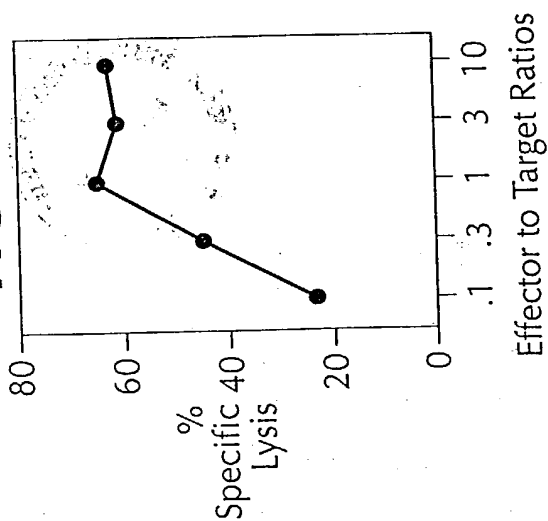
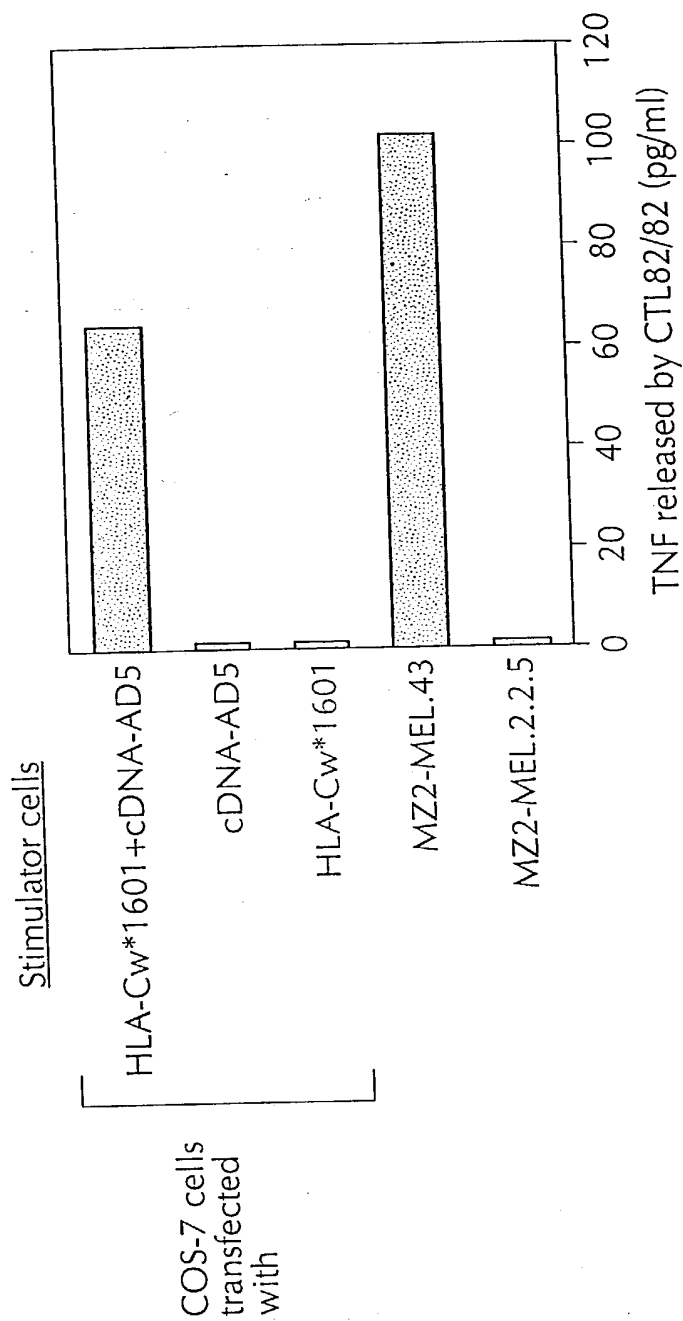


FIG. 1C



10001100 .040400

FIG. 2



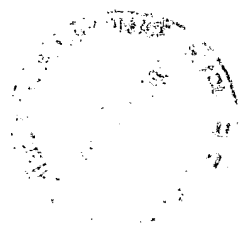


FIG. 3A

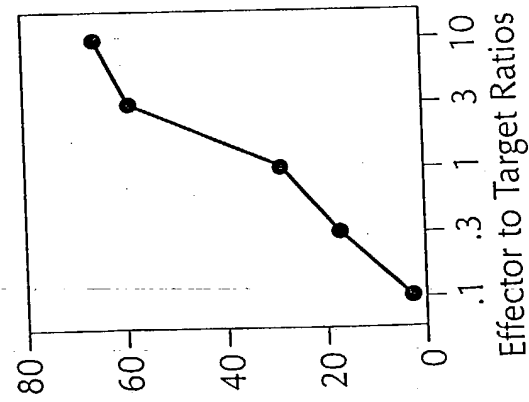


FIG. 3B

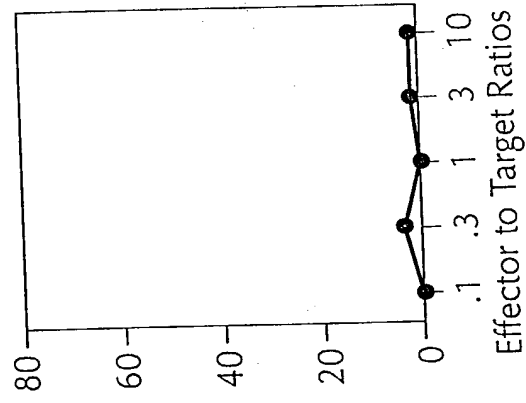


FIG. 3C

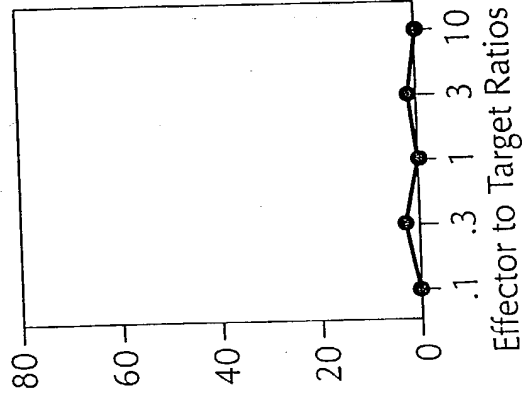


FIG. 3D

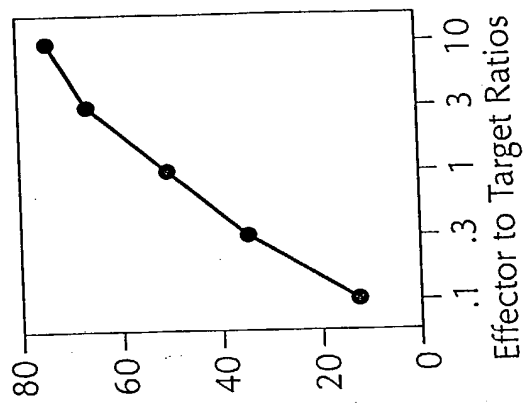


FIG. 4

CGCCAATTTA GGGTCTCCGG TATCTCCCGC TGAGCTGCTC TGTTCCCGGC TTAGAGGACC 60
 AGGAGAAAGG GGAGCTGGAG GCTGGAGCCT GTAAACACCGT GGCTCGTCTC ACTCTGGATG 120
 GTGGTGGCAA CAGAGATGGC AGCGCAGCTG GAGTGTTAGG AGGCGGCGCT GAGCGGTAGG 180
 AGTGGGGCTG GAGCAGTAAG ATGGCGGCCA GAGCGGTTTT TCTGGCATTG S A Q L
 L L Q A R L M K E E S P V V S W R L E P 240
 TGCTCCAAGC CAGGCTGATG AAGGAGGAGT CCCCTGTGGT GAGCTGGAGG TTGGAGCCTG 300
 E D G T A L C F I F 43
 AAGACGGCAC AGCTCTGTGC TTCATCTTCT GAGGTTGTGG CAGCCACGGT GATGGAGACG 360
 GCAGCTCAAC AGGAGCAATA GGAGGAGATG GAGTTTCACT GTGTCAGCCA GGATGGTCTC 420
 GATCTCCTGA CCTCGTGATC CGCCCGCCTT GGCCTTCCAA AGTGCCGAGA TTACAGCGAT 480
 GTGCATTTTG TAAGCACTTT GGAGCCACTA TCAAAATGCTG TGAAGAGAAA TGTACCCAGA 540
 TGTATCATTA TCCTTGTGCT GCAGGAGCCG GCTCCTTTCA GGATTTCACT CACATCTTCC 600
 TGCTTTGTCC AGAACACATT GACCAAGCTC CTGAAAGATG TAAAGTTTACT ACGCATAGAC 660
 TTTTAAACTT CAACCAATGT ATTTACTGAA AATAACAAAT GTTGTAATTT CCTGTAGTGT 730
 TATTCTACTT GTATTAAAAG GTAATAATAC ATAATCATTA AAATCTGAGG GATCATTTGCC 780
 AGAGATTGTT GGGGAGGGAA ATGTTATCAA CGGTTTCACT GAAATTAAT CCAAAAAGTT 840
 ATTTCCCTCAG AAAAATCAAA TAAAGTTTGC ATGTTTTTTA TTCTTAAAC ATTTTAAAAA 900
 CCACTGTAGA ATGATGTAAA TAGGGACTGT GCAGTATTTT TGACATATAC TATAAAATTA 960
 TTAAAAAGTC AATCAGTATT CAACATCTTT TACACTAAAA AGCC 1004

1000 1100 1200 1300 1400 1500

FIG. 5

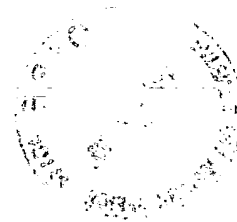
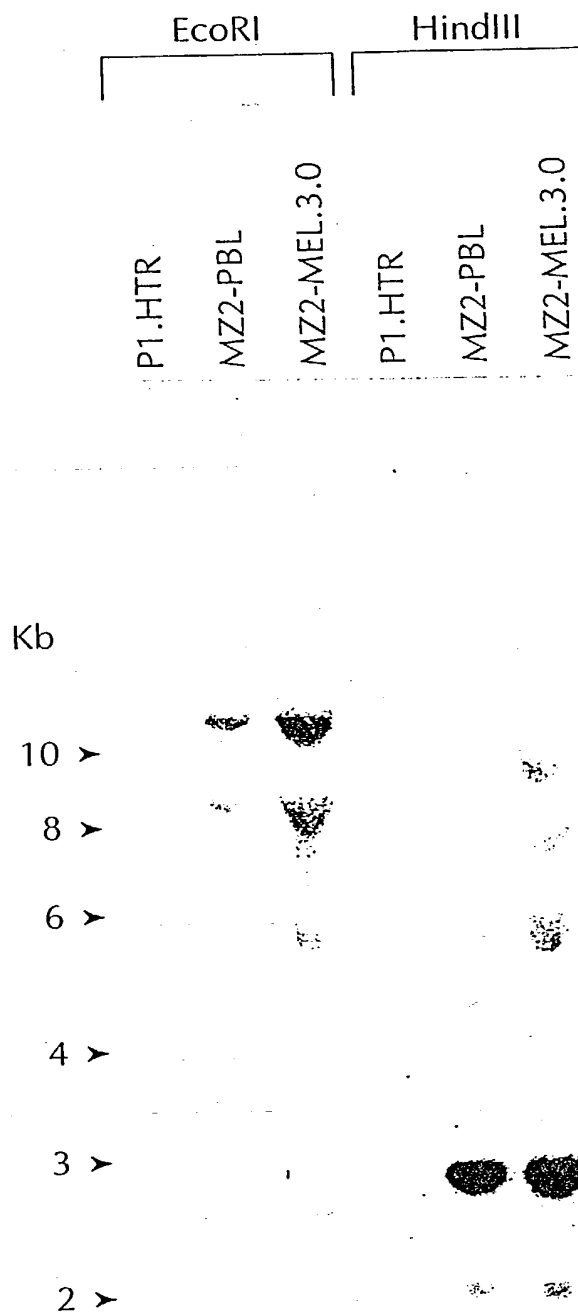


FIG. 6

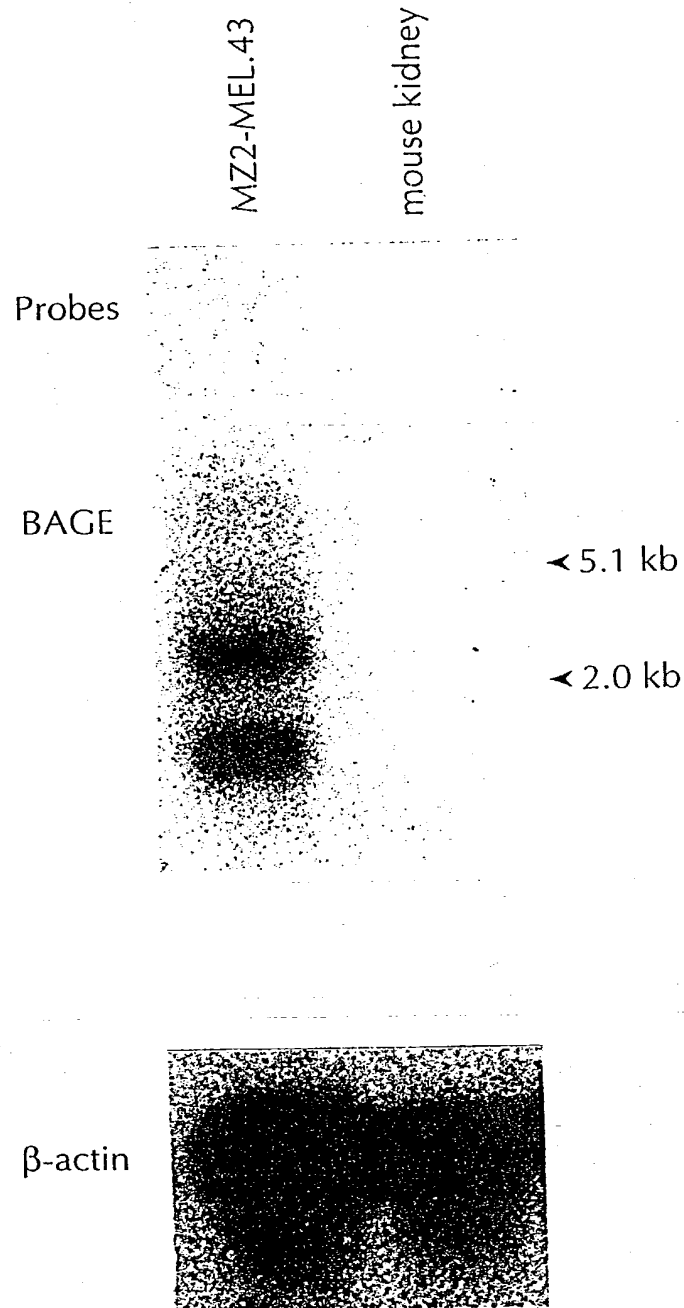


FIG. 7

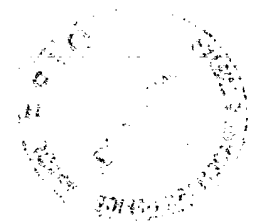
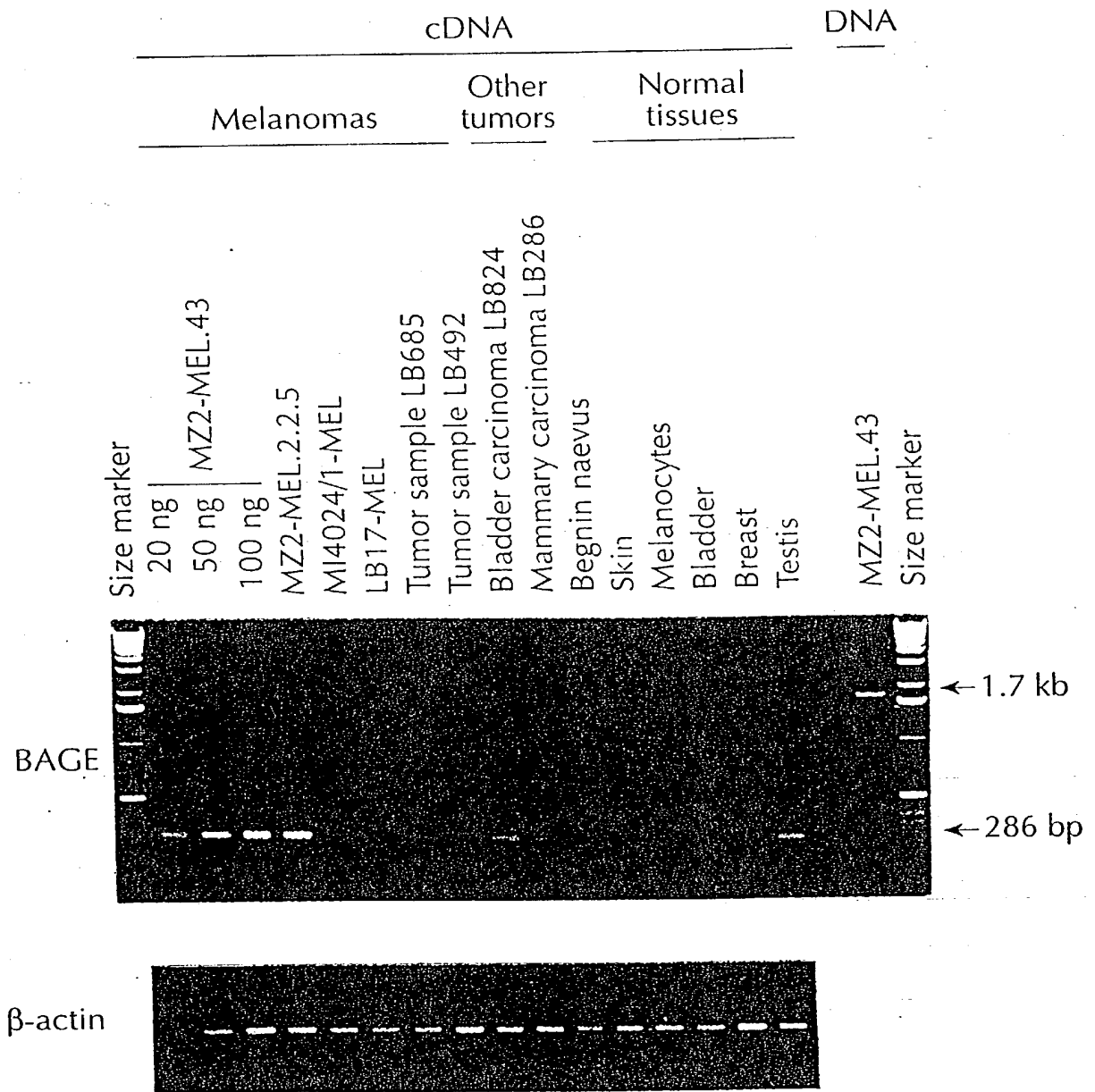


FIG. 8

